

US EPA  
Hazardous Waste Support Branch  
Validating Semivolatile Organic Compounds  
By Gas Chromatography/Mass Spectrometry  
SW-846 Method 8270D



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Annual Review

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YES NO N/A

## INTRODUCTION

## Scope and Applicability

This SOP offers detailed guidance in evaluating laboratory data generated according to "SW846-Method 8270D" January 1998. Method 8270D is used to determine the concentration of semivolatile organic compounds in extracts prepared from many types of solid waste matrices, soils, air sampling media and water samples. The validation methods and actions discussed in this document are based on the requirements set forth in SW846 Method 8270D, Method 8000C and the "USEPA Contract Laboratory Program National Functional Guidelines for Organic Data Review," January 2005. This document covers technical problems specific to each fraction and sample matrix; however, situations may arise where data limitations must be assessed based on the reviewer's professional judgement.

## Summary of Method

To ensure a thorough evaluation of each result in a data case, the reviewer must complete the checklist within this SOP, answering specific questions while performing the prescribed "ACTIONS" in each section. Qualifiers (or flags) are applied to questionable or unusable results as instructed. The data qualifiers discussed in this document are defined on page 5.

The reviewer must prepare a detailed data assessment to be submitted along with the completed SOP checklist. The Data Assessment must list all data qualifications, reasons for qualifications, instances of missing data and contract non-compliance.

## Reviewer Qualifications

Data reviewers must possess a working knowledge of SW846 Analytical Methods and National Functional Guidelines mentioned above.

## DEFINITIONS

### Acronyms

BNA - base neutral acid(another name for Semi Volatiles)  
CLP - Contract Laboratory Program  
CRQL - Contract Required Quantitation Limit  
%D - percent difference  
DCB -decachlorobiphenyl  
DDD - dichlorodiphenyldichloroethane  
DDE - dichlorodiphenylethane  
DDT - dichlorodiphenyltrichloroethane  
DoC - Date of Collection  
GC - gas chromatography  
GC/ECD - gas chromatograph/electron capture detector  
GC/MS - gas chromatograph/mass spectrometer  
GPC - gel permeation chromatography  
IS - internal standard  
kg - kilogram  
µg - microgram  
MS - matrix spike  
MSD - matrix spike duplicate  
ℓ - liter  
ml - milliliter  
PCB - Polychlorinated biphenyl  
PE - performance evaluation  
PEM - Performance Evaluation Mixture  
QC - quality control  
RAS - Routine Analytical Services  
RIC - reconstructed ion chromatogram  
RPD - relative percent difference  
RRF - relative response factor  
RRF - average relative response factor (from initial calibration)  
RRT - relative retention time  
RSD - relative standard deviation  
RT - retention time  
RSCC - Regional Sample Control Center  
SDG - sample delivery group  
SMC - system monitoring compound  
SOP - standard operating procedure  
SOW - Statement of Work  
SVOA - semivolatile organic acid  
TCL - Target Compound List  
TCLP - Toxicity Characteristics Leachate Procedure

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YES NO N/A

TCX -tetrachloro-m-xylene

TIC - tentatively identified compound

TOP0 - Task Order Project Officer

TPO - Technical Project Officer

VOA - Volatile organic

VTSR - Validated Time of Sample Receipt

## Data Qualifiers

- |    |   |   |
|----|---|---|
| U  | - | The analyte was analyzed for, but was not detected above the reported sample quantitation limit.  |
| J  | - | The analyte was positively identified; the associated numerical value is the approximate concentration of the analyte in the sample.  |
| N  | - | The analysis indicates the presence of an analyte for which there is presumptive evidence to make a "tentative identification."   |
| JN | - | The analysis indicates the presence of an analyte that has been "tentatively identified" and the associated numerical value represents its approximate concentration.   |
| UJ | - | The analyte was not detected above the reported sample quantitation limit. However, the reported quantitation limit is approximate and may or may not represent the actual limit of quantitation necessary to accurately and precisely measure the analyte in the sample. |
| R  | - | The sample results are rejected due to serious deficiencies in the ability to analyze the sample and meet quality control criteria. The presence or absence of the analyte cannot be verified.  |

**LAB QUALIFIERS:**

- |   |   |  |
|---|---|--|
| D | - | The positive value is the result of an analysis at a secondary dilution factor.  |
| B | - | The analyte is present in the associated method blank as well as in the sample. This qualifier has a different meaning when validating inorganic data. |

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YES NO N/A

- |        |   |   |
|--------|---|---|
| E      | - | The concentration of this analyte exceeds the calibration range of the instrument.  |
| A      | - | Indicates a Tentatively Identified Compound (TIC) is a suspected adol-condensation product.   |
| X,Y,Z- |   | Laboratory defined flags. The data reviewer must change these qualifiers during validation so that the data user may understand their impact on the data. |

## I. PACKAGE COMPLETENESS AND DELIVERABLES

CASE NUMBER: \_\_\_\_\_ LAB: \_\_\_\_\_

SITE NAME: \_\_\_\_\_

## 1.0 Data Completeness and Deliverables

1.1 Has all data been submitted in CLP deliverable format? [ ]

ACTION: If not, note the effect on review of the data in the data assessment narrative.

## 2.0 Cover Letter, SDG Narrative

2.1 Is a laboratory narrative or cover letter present? ☐ ☐ ☐

2.2 Are case number and SDG number(s) contained in the narrative or cover letter? ☐ ☐ ☐

## SEMIVOLATILE ANALYSES

## 1.0 Traffic Reports and Laboratory Narrative

\_\_\_\_\_

\_\_\_\_\_ [ ] \_\_\_\_\_

ACTION: If samples were not iced, or if the ice was melted upon arrival at the laboratory and the cooler temperature was elevated (10°C), flag all positive results "J" and all non-detects "UJ".

## 2.0 Holding Times

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40 days of the date of extraction.

Table of Holding Time Violations

(See Traffic Report)					
Sample ID	Sample Matrix	Date Sampled	Date Lab Received	Date Extracted	Date Analyzed
_____	_____	_____	_____	_____	_____
_____	_____	_____	_____	_____	_____
_____	_____	_____	_____	_____	_____
_____	_____	_____	_____	_____	_____
_____	_____	_____	_____	_____	_____
_____	_____	_____	_____	_____	_____

ACTION: If technical holding times are exceeded, flag all positive results as estimated ("J") and sample quantitation limits as estimated ("UJ"), and document in the narrative that holding times were exceeded.

If analyses were done more than 14 days beyond holding time, either on the first analysis or upon re analysis, the reviewer must use professional judgement to determine the reliability of the data and the effects of additional storage on the sample results. At a minimum, all results should be qualified "J", but the reviewer may determine that non-detect data are unusable ("R"). If holding times are exceeded by more than 28 days, all non-detect data are unusable (R).



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YES NO N/A

### 3.0 Surrogate Recovery (Form II/Equivalent)

3.1 Have the semi volatile surrogate recoveries been listed on CLP Surrogate Recovery forms (Form II) for each of the following matrices:

a. Low Water [ ]

b. Low/Med Soil [ ]

3.2 If so, are all the samples listed on the appropriate Surrogate Recovery Summary forms for each matrix:

a. Low Water [ ]

b. Low/Med Soil [ ]

ACTION: If CLP deliverables are unavailable, document the effect(s) in data assessments. In some cases the lab may have to be contacted to obtain the data necessary to complete the validation.

3.3 Were outliers marked correctly with an asterisk? [ ]

ACTION: Circle all outliers in red.

3.4 Were two or more base neutral OR acid surrogate recoveries out of specification for any sample or method blank (Reviewer should use lab in house recovery limits. Use surrogate recovery limits from USEPA National Functional Guidelines January 2005 page 130, if in house limits are not available. See Method 8000B-43 or 8000C-24). [ ]

Note: Examine lab in house limits for reasonableness.

If yes, were samples re-analyzed? [ ]

\_\_\_\_\_ [ ] \_\_\_\_\_

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effect in data assessments.

#### 4.0 Matrix Spikes (Form III/Equivalent)

4.1 Have the semivolatile Matrix Spike and Matrix Spike Duplicate/or duplicate unspiked Sample recoveries been listed on the Recovery Form (Form III)?

                          

NOTE: Method 3500B/page 4 states the spiking compounds:

Base/neutrals

1,2,4-Trichlorobenzene

Acenaphthene

2,4-Dinitrotoluene

Pyrene

N-Nitroso-di-n-propylamine

1,4-Dichlorobenzene

## Acids

## Pentachlorophenol

Phenol

2-Chlorophenol

4-Chloro-3-methylphenol

4-Nitrophenol

Note: Some projects may require the spiking of specific compounds of interest.

Note: See Method 8270D-sec 8.4.2 for deciding on whether to prepare and analyze duplicate samples or a matrix spike/matrix spike duplicate. If samples are expected to contain target analytes, then laboratory may use one matrix spike and a duplicate analysis of an unspiked field sample. If samples are not expected to contain target analytes, laboratory should use a matrix spike and matrix spike duplicate pair.

4.2 Were matrix spikes analyzed at the required frequency for each of the following matrices:

a. Low Water

[ ]

b. Low Solid

[ ]

c. Med Solid

[ ]

ACTION: If any matrix spike data are missing, take the action specified in 3.2 above. It may be necessary to contact the lab to obtain the required data.

NOTE: If the data has not been reported on CLP equivalent form, then the laboratory must provide the information necessary to evaluate the spike recoveries in the MS and MSD. The required data which should have been provided by the lab include the analytes and concentrations used for spiking, background concentrations of the spiked analytes (i.e., concentrations in unspiked sample), methods and equations used to calculate the QC acceptance criteria for the spiked analytes, percent recovery data for all spiked analytes.

The data reviewer must verify that all reported equations and percent recoveries are correct before proceeding to the next section.

4.3 Were matrix spikes performed at concentration equal to 100ug/L for acid compounds, and 200ug/l for base compounds (Method 3500B-4), or those specified in project plan. ☐ ☐ ☐

4.4 How many semivolatile spike recoveries are outside Laboratory in house MS/MSD recovery limits (use recovery limits values in Method 8270D-43&44 Table 6 if in house values not available).

Water  
  
\_\_\_\_ out of \_\_\_\_

Solids  
  
\_\_\_\_ out of \_\_\_\_

YES NO N/A

\_\_\_\_\_ out of \_\_\_\_\_

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YES NO N/A

the calibration standard or at any other time  
during the analytical shift for each GC/MS system  
used ?

☐ ☐ ☐

ACTION: If any method blank data are missing, call  
lab for explanation/resubmittal. If not  
available, use professional judgement to  
determine if the associated sample data  
should be qualified.

5.4 Chromatography: review the blank raw data -  
chromatograms (RICs), quant reports or data system  
printouts and spectra.

Is the chromatographic performance (baseline  
stability) for each instrument acceptable for  
the semivolatiles?

☐ ☐ ☐

ACTION: Use professional judgement to determine the  
effect on the data.

## 6.0 Contamination

NOTE: "Water blanks", "drill blanks" and "distilled  
water blanks" are validated like any other  
sample and are not used to qualify the data.  
Do not confuse them with the other QC blanks  
discussed below.

6.1 Do any method/instrument/reagent blanks have  
positive results for target analytes and/or TICs?  
When applied as described below, the contaminant  
concentration in these blanks are multiplied by  
the sample dilution factor and corrected for  
percent moisture where necessary.

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6.2 Do any field/rinse/ blanks have positive results  
for target analytes and/or TICs (if required,  
see section 10 below)?

☐ ☐ ☐

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YES NO N/A

ACTION: Prepare a list of the samples associated with each of the contaminated blanks.  
(Attach a separate sheet.)

NOTE: All field blank results associated to a particular group of samples (may exceed one per case) must be used to qualify data. Blanks may not be qualified because of contamination in another blank. Field Blanks must be qualified for outlying surrogates, poor spectra, instrument performance or calibration QC problems.

ACTION: Follow the directions in the table below to qualify sample results due to contamination. Use the largest value from all the associated blanks. If gross contamination exists, all data in the associated samples should be qualified as unusable (R).

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YES NO N/A

### Blank Action for Semivolatile Analyses

Blank Type	Blank Result	Sample Result	Action for Samples
Method, Field	Detects	Not detected	No qualification required
	< CRQL *	< CRQL	Report CRQL value with a U
		≥ CRQL	No qualification required
	= CRQL *	< CRQL	Report CRQL value with a U
		≥ CRQL	No qualification required
	> CRQL *	< CRQL	Report CRQL value with a U
		≥ CRQL and < blank contamination	Report concentration of sample with a U
		≥ CRQL and ≥ blank contamination	No qualification required

NOTE: Analytes qualified "U" for blank contamination are still considered as "hits" when qualifying for calibration criteria.

NOTE: If the laboratory did not report TIC analyses, check the project plans to verify whether or not it was required.

6.3 Are there field/rinse/equipment blanks associated with every sample?

ACTION: For low level samples, note in data assessment that there is no associated field/rinse/equipment blank. Exception: samples taken from a drinking water tap do not have associated field blanks.

6.4 Was a instrument blank analyzed after each sample/dilution which contained a target compound



YES NO N/A

that exceeded the initial calibration range. ☐ ☐ ☐

6.5 Does the instrument blank have positive results  
for target analytes and/or TICs? ☐ ☐ ☐

Note: Use professional judgement to determine  
if carryover occurred and qualify analytes  
accordingly.

#### 7.0 GC/MS Apparatus and Materials

7.1 Did the lab use the proper gas chromatographic  
column for analysis of semivolatiles by Method  
8270D? Check raw data, instrument logs or contact  
the lab to determine what type of column was used.  
The method requires the use of 30 m x 0.25 mm ID  
(or 0.32 mm ID), silicone-coated, fused silica,  
capillary column. ☐ ☐ ☐

ACTION: If the specified column, or equivalent, was  
not used, document the effects in the data  
assessment. Use professional judgement to  
determine the acceptability of the data.

#### 8.0 GC/MS Instrument Performance Check (Form V/Equivalent)

8.1 Are the GC/MS Instrument Performance Check Forms  
(Form V) present for decafluorotriphenylphosphine  
(DFTPP)? ☐ ☐ ☐

NOTE: The performance solution should also contain 4,4-DDT,  
pentachlorophenol, and benzidine to verify  
injection port inertness and column performance.  
The degradation of DDT to DDE and DDD must be  
less than 20% total and the response of  
pentachlorophenol and benzidine should be  
within normal ranges for these compounds (based  
upon lab experience) and show no peak degradation  
or tailing before samples are analyzed. (see section 5.5

page 8270D-12).

8.2 Are the enhanced bar graph spectrum and mass/charge (m/z) listing for the DFTPP provided for each twelve hour shift? ☐ ☐ ☐

8.3 Has an instrument performance check solution been analyzed for every twelve hours of sample analysis per instrument? ☐ ☐ ☐

ACTION: List date, time, instrument ID, and sample analyses for which no associated GC/MS tuning data are available.

DATE	TIME	INSTRUMENT	SAMPLE NUMBERS
_____	_____	_____	_____
_____	_____	_____	_____
_____	_____	_____	_____

ACTION: If lab cannot provide missing data, reject ("R") all data generated outside an acceptable twelve hour calibration interval.

ACTION: If mass assignment is in error, flag all associated sample data as unusable (R).

8.4 Have the ion abundances been normalized to m/z 198? ☐ ☐ ☐

8.5 Have the ion abundance criteria been met for each instrument used? ☐ ☐ ☐

ACTION: List all data which do not meet ion abundance criteria (attach a separate sheet).

ACTION: If ion abundance criteria are not met, take  
action specified in section 3.2

8.6 Are there any transcription/calculation errors  
between mass lists and Form Vs? (Check at least  
two values but if errors are found, check more.) ☐ ☐ ☐

8.7 Have the appropriate number of significant  
figures (two) been reported? ☐ ☐ ☐

ACTION: If large errors exist, call lab for  
explanation/resubmittal, make necessary  
corrections and document effect in data  
assessments.

8.8 Are the spectra of the mass calibration compound  
acceptable? ☐ ☐ ☐

ACTION: Use professional judgement to determine  
whether associated data should be accepted,  
qualified, or rejected.

## 9.0 Target Analytes

9.1 Are the Organic Analysis Data Sheets (Form I)  
present with required header information on each  
page, for each of the following:

a. Samples and/or fractions as appropriate ☐ ☐ ☐

b. Matrix spikes and matrix spike duplicates ☐ ☐ ☐

c. Blanks ☐ ☐ ☐

9.2 Has any special cleanup, such as GPC, been  
performed on all soil/sediment sample extracts  
(see section 7.2, page 8270D-14)? ☐ ☐ ☐

ACTION: If data suggests that extract cleanup was not performed, use professional judgement. Make note in the data assessment narrative.

9.3 Are the Reconstructed Ion Chromatograms, mass spectra for the identified compounds, and the data system printouts (Quant Reports) included in the sample package for each of the following?

- |   |                          |     |     |
|---|--------------------------|-----|-----|
| a. Samples and/or fractions as appropriate                                  | <input type="checkbox"/> | ___ | ___ |
| b. Matrix spikes and matrix spike duplicates<br>(Mass spectra not required) | <input type="checkbox"/> | ___ | ___ |
| c. Blanks   | <input type="checkbox"/> | ___ | ___ |

ACTION: If any data are missing, take action specified in 3.2 above.

9.4 Are the response factors shown in the Quant Report? ☐ \_\_\_ \_\_\_

9.5 Is chromatographic performance acceptable with respect to:

- |                                 |                          |     |     |
|---------------------------------|--------------------------|-----|-----|
| Baseline stability?             | <input type="checkbox"/> | ___ | ___ |
| Resolution?                     | <input type="checkbox"/> | ___ | ___ |
| Peak shape?                     | <input type="checkbox"/> | ___ | ___ |
| Full-scale graph (attenuation)? | <input type="checkbox"/> | ___ | ___ |
| Other:_____                     | <input type="checkbox"/> | ___ | ___ |

ACTION: Use professional judgement to determine the acceptability of the data.

9.6 Are the lab-generated standard mass spectra of identified semivolatile compounds present for

YES NO N/A

each sample? ☐ ☐ ☐

ACTION: If any mass spectra are missing, take action specified in 3.2 above. If the lab does not generate their own standard spectra, make a note in the data assessment narrative. If spectra are missing, reject all positive data.

9.7 Is the RRT of each reported compound within 0.06 RRT units of the standard RRT in the continuing calibration? ☐ ☐ ☐

9.8 Are all ions present in the standard mass spectrum at a relative intensity greater than 10% (of the most abundant ion) also present in the sample mass spectrum? ☐ ☐ ☐

9.9 Do the relative intensities of the characteristic ions in the sample agree within  $\pm 30\%$  of the corresponding relative intensities in the reference spectrum? ☐ ☐ ☐

ACTION: Use professional judgement to determine acceptability of data. If it is determined that incorrect identifications were made, all such data should be rejected (R), flagged "N" (Presumptive evidence of the presence of the compound) or changed to not detected (U) at the calculated detection limit. In order to be positively identified, the data must comply with the criteria listed in 9.7, 9.8, and 9.9.

ACTION: When sample carry-over is a possibility, professional judgement should be used to determine if instrument cross-contamination has affected any positive compound identification.

10.0 Tentatively Identified Compounds (TIC)

10.1 If Tentatively Identified Compounds were required for this project, are all Form Is, Part B present; and do listed TICs include scan number or retention time, estimated concentration and "JN" qualifier?

NOTE: Review sampling reports to determine if the lab was required to identify non target analytes (refer to section 7.6.2,page 8270D-21).

10.2 Are the mass spectra for the tentatively identified compounds and associated "best match" spectra included in the sample package for each ☐ \_\_\_ of the following:

a. Samples and/or fractions as appropriate ☐ \_\_\_

b. Blanks ☐ \_\_\_

ACTION: If any TIC data are missing, take action specified in 3.2 above.

ACTION: Add "JN" qualifier only to analytes identified by CAS #.

10.3 Are any target compounds from one fraction listed as TIC compounds in another (e.g., an acid compound listed as a base neutral TIC)? \_\_\_ ☐ \_\_\_

ACTION: i. Flag with "R" any target compound listed as a TIC.

ii. Make sure all rejected compounds are properly reported in the other fraction.

10.4 Are all ions present in the reference mass spectrum with a relative intensity greater than 10% (of the most abundant ion) also present in the

YES NO N/A

sample mass spectrum? ☐ ☐ ☐

10.5 Do TIC and "best match" standard relative ion intensities agree within  $\pm 20\%$ ? ☐ ☐ ☐

ACTION: Use professional judgement to determine acceptability of TIC identifications. If it is determined that an incorrect identification was made, change the identification to "unknown" or to some less specific identification (example: "C3 substituted benzene") as appropriate and remove "JN". Also, when a compound is not found in any blank, but is a suspected artifact of a common laboratory contaminant, the result should be qualified as unusable, "R."

#### 11.0 Compound Quantitation and Reported Detection Limits

11.1 Are there any transcription/calculation errors in Form I results? Check at least two positive values. Verify that the correct internal standard, quantitation ion, and RRF were used to calculate Form I result. Were any errors found? ☐ ☐ ☐

NOTE: Structural isomers with similar mass spectra, but insufficient GC resolution (i.e. percent valley between the two peaks  $> 25\%$ ) should be reported as isomeric pairs. The reviewer should check the raw data to ensure that all such isomers were included in the quantitation (i.e., add the areas of the two coeluting peaks to calculate the total concentration).

11.2 Are the method detection limits adjusted to reflect sample dilutions and, for soils, sample moisture? ☐ ☐ ☐

ACTION: If errors are large, call lab for explanation/resubmittal, make any necessary corrections and document effect in data assessments.

ACTION: When a sample is analyzed at more than one dilution, the lowest detection limits are used (unless a QC exceedance dictates the use of the higher detection limit from the diluted sample data). Replace concentrations that exceed the calibration range in the original analysis by crossing out the "E" and it's associated value on the original Form I (if present) and substituting the data from the analysis of the diluted sample. Specify which Form I is to be used, then draw a red "X" across the entire page of all Form I's that should not be used, including any in the summary package.

12.0 Standards Data (GC/MS)

12.1 Are the Reconstructed Ion Chromatograms, and data system printouts (Quant, Reports) present for initial and continuing calibration? ☐ ☐ ☐

ACTION: If any calibration standard data are missing, take action specified in 3.2 above.

13.0 GC/MS Initial Calibration (Form VI/Equivalent)

13.1 Is the Initial Calibration Form (Form VI/Equivalent) present and complete for the semivolatile fraction? ☐ ☐ ☐

ACTION: If any calibration forms or standard row data are missing, take action specified in 3.2 above.

13.2 Are all base neutral or acid RRFs > 0.050? ☐ ☐ ☐



Check the **average RRFs** of the four System Performance Check Compounds (SPCCs):  
N-nitroso-di-n-propylamine, hexachlorocyclopentadiene, 2,4-dinitrophenol, and 4-nitrophenol. These compounds must have **average RRFs** greater than or equal to 0.05 before running samples and should not show any peak tailing.

ACTION: Circle all outliers in red.

ACTION: For any target analyte with **average RRF <0.05**

1. "R" all non-detects;
2. "J" all positive results.

13.3 Are response factors for base neutral or acid target analytes stable over the concentration range of the calibration (% Relative standard deviation [%RSD] < 15.0%)? [ ] \_ \_

NOTE: The % RSD for each individual Calibration Check Compound (CCC, Method 8270D-40 see Table 4) must be less than 30% before analysis can begin. If greater 30%, the lab must clean and recalibrate the instrument.

#### CALIBRATION CHECK COMPOUNDS

Base/Neutral Fraction	Acid Fraction
Acenaphthene	4-Chloro-3-methylphenol
1,4-Dichlorobenzene	2,4-Dichlorophenol
Hexachlorobutadiene	2-Nitrophenol
Diphenylamine	Phenol
Di-n-octyl phthalate	Pentachlorophenol
Fluoranthene	2,4,6-Trichlorophenol

YES NO N/A

Benzo ( a ) pyrene

ACTION: If the %RSD for any CCC >30% and no corrective action taken, then "J" qualify all positive hits and "UJ" qualify all non-detects.

ACTION: Circle all outliers in red.

ACTION: If the % RSD is  $\geq 15.0\%$ , qualify positive results for that analyte "J" and non-detects using professional judgement. When  $RSD > 90\%$ , flag all non-detect results for that analyte "R," unusable. Alternatively, the lab should calculate first or second order regression fit of the calibration curve and select the fit which introduces the least amount of error.

NOTE: Analytes previously qualified "U" due to blank contamination are still considered as "hits" when qualifying for calibration criteria.

13.4 Did the laboratory calculate the calibration curve by the least squares regression fit?

13.5 Are there any transcription/calculation errors in the reporting of average response factors (RRF) or % RSD? (Check at least two values but if errors are found, check more.)

ACTION: Circle Errors in red.

ACTION: If errors are large, call lab for explanation/resubmittal, make any necessary corrections and note errors in data assessments.

13.5 Do the target compounds for this SDG include Pesticides?

13.6 If the pesticide compounds include DDT, was the percent breakdown of DDT to DDD and DDE greater than 20%? ☐ ☐ ☐

ACTION: If DDT percent breakdown exceeds 20%:

- i. Qualify all positive results for DDT with "J". If DDT was not detected, but DDD and DDE results are positive, qualify the quantitation limit for DDT as unusable, "R".
- ii. Qualify all positive results for DDD and DDE as presumptively present at an approximate concentration "JN".

14.0 GC/MS Calibration Verification (Form VII/Equivalent)

14.1 Are the Calibration Verification Forms (Form VII) present and complete for all compounds of interest? ☐ ☐ ☐

14.2 Has a calibration verification standard been analyzed for every twelve hours of sample analysis per instrument? ☐ ☐ ☐

ACTION: List below all sample analyses that were not within twelve hours of a calibration verification analysis for each instrument used.

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ACTION: If any forms are missing or no calibration verification standard has been analyzed within twelve hours of every sample analysis,

YES NO N/A

call lab for explanation/resubmittal. If continuing calibration data are not available, flag all associated sample data as unusable ("R").

14.3 Do any of the SPCCs have an RRF <0.05? ☐ ☐ ☐

If YES, make a note in data assessment if the lab did not take corrective action specified in section 7.4.4, page 8270D-18. ☐ ☐ ☐

14.4 Do any of the CCCs have a %D between the initial and continuing RRF which exceeds 20.0%?

ACTION: If yes, make a note in data assessment.

14.5 Do any semivolatile compounds have a % Difference (% D) between the initial and continuing RRF which exceeds 20.0%? ☐ ☐ ☐

ACTION: Circle all outliers in red.

ACTION: Qualify both positive results and non-detects for the outlier compound(s) as estimated (J). When %D is above 90%, qualify all non-detects for that analyte as "R", unusable.

14.6 Do any semivolatile compounds have a RRF < 0.05? ☐ ☐ ☐

ACTION: Circle all outliers in red.

ACTION: If RRF < 0.05, qualify as unusable ("R") associated non-detects and "J" associated positive values.

14.7 Are there any transcription/calculation errors in the reporting of average response factors (RRF) or percent difference (%D) between initial and continuing RRFs? (Check at least two values but if errors are found, check more). ☐ ☐ ☐

ACTION: Circle errors in red.

ACTION: If errors are large, call lab for explanation/resubmittal, make any necessary corrections and document effect(s) in the data assessments.

15.0 Internal Standards (Form VIII)

15.1 Are the internal standard areas (Form VIII) of every sample and blank within the upper and lower limits (-50% to + 100%) for each continuing calibration? ☐ ☐ ☐

ACTION: List each outlying internal standard below.

Sample ID	IS #	Area	LowerLimit	Upper Limit
_____	_____	_____	_____	_____
_____	_____	_____	_____	_____
_____	_____	_____	_____	_____
_____	_____	_____	_____	_____

(Attach additional sheets if necessary.)

Note: Check Table 5, 8270D-41 for associated analytes.

- ACTION:
- If the internal standard area count is outside the upper or lower limit, flag with "J" all positive results and non-detects (U values) quantitated with this internal standard.
  - Non-detects associated with IS > 100% should not be qualified.

S)))))))))))))S)))))))))))))Q

YES NO N/A

- iii. If the IS area is below the lower limit (<50%), qualify all associated non-detects (U-values) "J". If extremely low area counts are reported (<25%) or if performance exhibits a major abrupt drop off, flag all associated non-detects as unusable (R).

15.2 Are the retention times of all internal standards within 30 seconds of the associated calibration standard?

                          

ACTION: Professional judgement should be used to qualify data if the retention times differ by more than 30 seconds.

## 16.0 Laboratory Control Samples (LCS)

16.1 Were any LCS samples run in order to verify analytes which failed criteria for spike recovery?

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16.2 Did the lab spike LCS sample spiked with the same analytes and the same concentrations as the matrix spike?

                          

16.3 Were the mean and standard deviation of all analytes within the QC acceptance ranges as shown in Table 6, 8270D-43?

[ ]      \_\_\_\_\_

ACTION: If the recovery of any analyte falls out of the designated range, the analytical results for that compound is suspect and should be qualified "J" in the unspiked samples.

## 17.0 Field Duplicates

17.1 Were any field duplicates submitted for semivolatile analysis?

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S))Q

YES NO N/A

ACTION: Compare the reported results for field duplicates and calculate the relative percent difference.

ACTION: Any gross variation between field duplicate results must be addressed in the reviewer narrative. However, if large differences exist, identification of field duplicates should be confirmed by contacting the sampler.